PCT09

```
RAW SEQUENCE LISTING                        DATE: 01/22/2002
PATENT APPLICATION:  US/09/700,130          TIME: 10:21:15

Input Set : A:\SEQLISTPLOUG1.001APC.txt
Output Set: N:\CRF3\01182002\I700130.raw
```

Does Not Comply
Corrected Diskette Needed

```
 4 <110> APPLICANT: Gerdes, Kenn
 5        Mikkelson, Marie (FKA Gotfredsen)
 6        Gronlund, Hugo
 7        Pedersen, Kim
 8        Kristoffersen, Peter
10 <120> TITLE OF INVENTION: CYTOTOXIN-BASED BIOLOGICAL CONTAINMENT
13 <130> FILE REFERENCE: PLOUG1.001APC
15 <140> CURRENT APPLICATION NUMBER: US 09/700,130
16 <141> CURRENT FILING DATE: 2000-11-07
18 <150> PRIOR APPLICATION NUMBER: PCT/DK99/00258
19 <151> PRIOR FILING DATE: 1999-05-07
21 <150> PRIOR APPLICATION NUMBER: DK 0627/98
22 <151> PRIOR FILING DATE: 1998-05-07
24 <150> PRIOR APPLICATION NUMBER: US 60/085,067
25 <151> PRIOR FILING DATE: 1998-05-12
27 <160> NUMBER OF SEQ ID NOS: 59
29 <170> SOFTWARE: FastSEQ for Windows Version 4.0
```

ERRORED SEQUENCES

```
1190 <210> SEQ ID NO: 59
1191 <211> LENGTH: 23
1192 <212> TYPE: DNA
1193 <213> ORGANISM: Artificial Sequence
1195 <220> FEATURE:
1196 <223> OTHER INFORMATION: primer AS-relE was used for the amplification of
1197       the relE coding region from the plasmid pMG223
1199 <400> SEQUENCE: 59
1200 gagacccac actaccatcg gcg                                    23
E--> 1201 ()
```

```
                    VERIFICATION SUMMARY              DATE: 01/22/2002
                    PATENT APPLICATION: US/09/700,130  TIME: 10:21:17

                    Input Set : A:\SEQLISTPLOUG1.001APC.txt
                    Output Set: N:\CRF3\01182002\I700130.raw

L:155 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:159 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:165 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:168 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:967 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:44
L:967 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:44
L:967 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:972 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:44
L:972 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:44
L:972 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:976 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:44
L:976 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:44
L:976 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:997 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:45
L:997 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:45
L:997 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:1003 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:45
L:1003 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:45
L:1003 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:1008 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:45
L:1008 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:45
L:1008 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:1031 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:46
L:1031 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:46
L:1031 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46
L:1035 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:46
L:1035 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:46
L:1035 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46
L:1201 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:23 SEQ:59
```